

SEQUENCE LISTING

ADEMAN 110>	Pang and Lu	
<120>	METHODS AND REAGENTS FOR TREATING GLUCOSE METABOLIC DISORDER	S
<130>	CIBT-P01-058	
<140> <141>	09/499526 2000-02-10	RECEIVED
	60/119,577 1999-02-10	JUL 0 9 2003
<160>	TEC	H CENTER 1600/2900
<170>	PatentIn version 3.1	
<210> <211> <212> <213>	1 582 DNA Homo sapiens	
<220> <221> <222> <223>	CDS (81)(371)	·
<400> cagctt	1 gace tgeggeagtg cagecettgg gactteeste geetteeace teetgetegt	60
ctgctt	Met Val Phe Val Arg Arg Pro Trp Pro Ala Leu 1 5 10	113
	a gtg ctt ctg gcc ctg ctc gtc tgc cta ggg gcg ctg gtc gac : Val Leu Leu Ala Leu Leu Val Cys Leu Gly Ala Leu Val Asp 15 20 25	161
	c ccc atc aaa ccc gag gct ccc ggc gaa gac gcc tcg ccg gag c Pro Ile Lys Pro Glu Ala Pro Gly Glu Asp Ala Ser Pro Glu 30 35 40	209
	g aac cgc tac tac gcc tcc ctg cgc cac tac ctc aac ctg gtc n Asn Arg Tyr Tyr Ala Ser Leu Arg His Tyr Leu Asn Leu Val 50 55	257
-	g cag cgg tat ggg aaa aga gac ggc ccg gac agg ctt ctt tcc g Gln Arg Tyr Gly Lys Arg Asp Gly Pro Asp Arg Leu Leu Ser 65 70 75	305
	g ttc ttc ccc gac ggc gag gac cgc ccc gtc agg tcg cgg tcg r Phe Phe Pro Asp Gly Glu Asp Arg Pro Val Arg Ser Arg Ser 80 85 90	353
	c cca gac ctg tgg tgaggacccc tgaggcctcc tgggagatct y Pro Asp Leu Trp 95	401
gccaac	cacg cccacgtcat ttgcatacgc actcccgacc ccagaaaccc ggattctgcc	461

521

581

582

teecgaegge ggegtetggg eagggttegg gtgeggeeet eeggeeggt eteggtgeee ccgcccctg ggctggaggg ctgtgtgtgg tccttccctg gtcccaaaat aaagagcaaa t <210> 2 <211> 97 <212> PRT <213> Homo sapiens <400> 2 Met Val Phe Val Arg Arg Pro Trp Pro Ala Leu Thr Thr Val Leu Leu Ala Leu Leu Val Cys Leu Gly Ala Leu Val Asp Alan Tyr Pro Ile Lys Pro Glu Ala Pro Gly Glu Asp Ala Ser Pro Glu Glu Leu Asn Arg Tyr 40 Tyr Ala Ser Leu Arg His Tyr Leu Asn Leu Val Thr Arg Gln Arg Tyr 55 60 Gly Lys Arg Asp Gly Pro Asp Arg Leu Leu Ser Lys Thr Phe Pro 70 75 Asp Gly Glu Asp Arg Pro Val Arg Ser Arg Ser Glu Gly Pro Asp Leu Trp <210> 3 <211> 36 <212> PRT <213> Homo sapiens <400> 3 Tyr Pro Ile Lys Pro Glu Ala Pro Gly Glu Asp Ala Ser Pro Glu Glu Leu Asn Arg Tyr Tyr Ala Ser Leu Arg His Tyr Leu Asn Leu Val Thr Arg Gln Arg Tyr

35